

SEQUENCE LISTING

<110> TOLEDANO, MICHEL
BITEAU, BENOIT

<120> APPLICATIONS OF A NEW CLASS OF ENZYMES: SULFIREDOXINS

<130> 1169-042

<140> 10/563,375

<141> 2006-01-04

<150> PCT/FR04/01727

<151> 2004-07-02

<150> FR 03/08212

<151> 2003-07-04

<160> 17

<170> PatentIn Ver. 3.3

<210> 1

<211> 127

<212> PRT

<213> *Saccharomyces cerevisiae*

<400> 1

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Ser | Leu | Gln | Ser | Asn | Ser | Val | Lys | Pro | Thr | Glu | Ile | Pro | Leu | Ser |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Glu | Ile | Arg | Arg | Pro | Leu | Ala | Pro | Val | Leu | Asp | Pro | Gln | Lys | Ile | Asp |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| Ala | Met | Val | Ala | Thr | Met | Lys | Gly | Ile | Pro | Thr | Ala | Ser | Lys | Thr | Cys |
| | | 35 | | | | | 40 | | | | | 45 | | | |
| Ser | Leu | Glu | Gln | Ala | Glu | Ala | Ala | Ala | Ser | Ala | Gly | Glu | Leu | Pro | Pro |
| | 50 | | | | | 55 | | | | | 60 | | | | |
| Val | Asp | Val | Leu | Gly | Val | Arg | Val | Lys | Gly | Gln | Thr | Leu | Tyr | Tyr | Ala |
| 65 | | | | | 70 | | | | | 75 | | | | | 80 |
| Phe | Gly | Gly | Cys | His | Arg | Leu | Gln | Ala | Tyr | Asp | Arg | Arg | Ala | Arg | Glu |
| | | | | 85 | | | | | 90 | | | | | 95 | |
| Thr | Gln | Asn | Ala | Ala | Phe | Pro | Val | Arg | Cys | Arg | Val | Leu | Pro | Ala | Thr |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| Pro | Arg | Gln | Ile | Arg | Met | Tyr | Leu | Gly | Ser | Ser | Leu | Asp | Ile | Glu | |
| | | 115 | | | | | 120 | | | | | 125 | | | |

<210> 2

<211> 120

<212> PRT

<213> *Candida albicans*

<400> 2

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Met Ser Met Tyr Thr Ser Arg Leu Ala Thr Glu Tyr Val Pro Leu Ser
 1              5              10              15

Glu Ile Lys Arg Pro Ile Pro Pro Val Leu Asp Tyr Gln Lys Ile Asp
      20              25              30

Ala Met Leu Ser Thr Leu Lys Gly Val Pro Met Glu Ser Ala Thr Cys
      35              40              45

Lys Val Glu Asp Ile Thr Ala Gly Glu Leu Pro Pro Ile Asp Val Phe
      50              55              60

Lys Ile Arg Glu Asn Gly Lys Asn Phe Tyr Phe Ala Phe Gly Gly Cys
      65              70              75              80

His Arg Phe Gln Ala Tyr Asp Arg Ile Ser Lys Glu Thr Glu Lys Glu
      85              90              95

Val Met Val Lys Ser Arg Ile Leu Pro Ala Thr Arg Lys Ser Leu Arg
      100              105              110

Ile Tyr Leu Gly Ala Ser Val Asp
      115              120

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<210> 3

<211> 124

<212> PRT

<213> Schizosaccharomyces pombe

<400> 3

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Met Thr Ser Ile His Thr Gly Ser Asn Asn Asn Ile Val Glu Leu Asp
 1              5              10              15

Met Ser Glu Leu Ile Arg Pro Ile Pro Pro Val Leu Asp Met Asn Lys
      20              25              30

Val Asn Ser Met Met Glu Thr Met Thr Gly Lys Thr Pro Pro Ala Ser
      35              40              45

Cys Gly Leu Thr Ser Glu Asp Leu Glu Ala Gly Glu Leu Pro Pro Val
      50              55              60

Asp Val Leu Thr Phe Lys Lys Ser Gly Lys Pro Tyr Tyr Phe Ala Phe
      65              70              75              80

Gly Gly Cys His Arg Leu Arg Ala His Asp Glu Ala Gly Arg Lys Lys
      85              90              95

Val Arg Cys Lys Leu Val Asn Cys Ser Pro Asn Thr Leu Arg Leu Tyr
      100              105              110

Leu Gly Ala Ser Ala Asn Lys Phe Leu Asp Ser Asp
      115              120

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<210> 4
 <211> 137
 <212> PRT
 <213> Homo sapiens

<400> 4
 Met Gly Leu Arg Ala Gly Gly Thr Leu Gly Arg Ala Gly Ala Gly Arg
 1 5 10 15
 Gly Ala Pro Glu Gly Pro Gly Pro Ser Gly Gly Ala Gln Gly Gly Ser
 20 25 30
 Ile His Ser Gly Arg Ile Ala Ala Val His Asn Val Pro Leu Ser Val
 35 40 45
 Leu Ile Arg Pro Leu Pro Ser Val Leu Asp Pro Ala Lys Val Gln Ser
 50 55 60
 Leu Val Asp Thr Ile Arg Glu Asp Pro Asp Ser Val Pro Pro Ile Asp
 65 70 75 80
 Val Leu Trp Ile Lys Gly Ala Gln Gly Gly Asp Tyr Phe Tyr Ser Phe
 85 90 95
 Gly Gly Cys His Arg Tyr Ala Ala Tyr Gln Gln Leu Gln Arg Glu Thr
 100 105 110
 Ile Pro Ala Lys Leu Val Gln Ser Thr Leu Ser Asp Leu Arg Val Tyr
 115 120 125
 Leu Gly Ala Ser Thr Pro Asp Leu Gln
 130 135

<210> 5
 <211> 136
 <212> PRT
 <213> Mus musculus

<400> 5
 Met Gly Leu Arg Ala Gly Gly Ala Leu Arg Arg Ala Gly Ala Gly Pro
 1 5 10 15
 Gly Ala Pro Val Val His Gly Pro Gly Gly Ala Gln Gly Gly Ser Ile
 20 25 30
 His Ser Gly Cys Ile Ala Thr Val His Asn Val Pro Ile Ala Val Leu
 35 40 45
 Ile Arg Pro Leu Pro Ser Val Leu Asp Pro Ala Lys Val Gln Ser Leu
 50 55 60
 Val Asp Thr Ile Leu Ala Asp Pro Asp Ser Val Pro Pro Ile Asp Val
 65 70 75 80
 Leu Trp Ile Lys Gly Ala Gln Gly Gly Asp Tyr Tyr Tyr Ser Phe Gly
 85 90 95

Gly Cys His Arg Tyr Ala Ala Tyr Gln Gln Leu Gln Arg Glu Thr Ile
 100 105 110

Pro Ala Lys Leu Val Arg Ser Thr Leu Ser Asp Leu Arg Met Tyr Leu
 115 120 125

Gly Ala Ser Thr Pro Asp Leu Gln
 130 135

<210> 6

<211> 162

<212> PRT

<213> *Drosophila melanogaster*

<400> 6

Met Glu Phe Ile Ser His Phe Leu Arg Ala Thr Ser Arg Arg Thr Ala
 1 5 10 15

Ala Leu Gly Pro Ile Leu Gln Arg Asn Arg Ser Glu Ile Ile Gln Lys
 20 25 30

Gln Ser Leu Thr Asn Arg Gln Ala Phe Arg Arg Tyr Arg Ser Ser Cys
 35 40 45

Ser Thr Met Asp Thr Thr Val His Ser Ala Gly Ile Asp Glu Thr His
 50 55 60

Leu Val Pro Met Ser Val Ile Gln Arg Pro Ile Pro Ser Val Leu Asp
 65 70 75 80

Glu Gln Lys Val Gln Ser Leu Met Glu Thr Ile Lys Asn Glu Thr Ser
 85 90 95

Glu Asp Glu Val Pro Pro Ile Asp Leu Leu Trp Ile Ser Gly Ser Glu
 100 105 110

Gly Gly Asp Tyr Tyr Phe Ser Phe Gly Gly Cys His Arg Phe Glu Ala
 115 120 125

Tyr Lys Arg Leu Gln Arg Pro Thr Ile Lys Ala Lys Leu Val Lys Ser
 130 135 140

Thr Leu Gly Asp Leu Tyr His Tyr Met Gly Ser Ser Ala Pro Lys Tyr
 145 150 155 160

Leu Ala

<210> 7

<211> 125

<212> PRT

<213> *Arabidopsis thaliana*

<400> 7

Met Ala Asn Leu Met Met Arg Leu Pro Ile Ser Leu Arg Ser Phe Ser
 1 5 10 15

Val Ser Ala Ser Ser Ser Asn Gly Ser Pro Pro Val Ile Gly Gly Ser
 20 25 30
 Ser Gly Gly Val Gly Pro Met Ile Val Glu Leu Pro Leu Glu Lys Ile
 35 40 45
 Arg Arg Pro Leu Met Arg Thr Arg Ser Asn Asp Gln Asn Lys Val Lys
 50 55 60
 Glu Leu Met Asp Ser Ile Arg Gln Ile Gly Leu Gln Val Pro Ile Asp
 65 70 75 80
 Val Ile Glu Val Asp Gly Thr Tyr Tyr Gly Phe Ser Gly Cys His Arg
 85 90 95
 Tyr Glu Ala His Gln Lys Leu Gly Leu Pro Thr Ile Arg Cys Lys Ile
 100 105 110
 Arg Lys Gly Thr Lys Glu Thr Leu Arg His His Leu Arg
 115 120 125

<210> 8
 <211> 86
 <212> PRT
 <213> Thermosynechococcus elongatus

<400> 8
 Met Arg Val Leu Asp Leu Pro Leu Asn Ala Ile Arg Arg Pro Leu Val
 1 5 10 15
 Arg Gln Thr Asp Pro Ala Lys Val Ala Ala Leu Met Ala Ser Ile Ala
 20 25 30
 Glu Ile Gly Gln Gln Glu Pro Ile Asp Val Leu Glu Val Glu Gly His
 35 40 45
 Tyr Tyr Gly Phe Ser Gly Cys His Arg Tyr Glu Ala Cys Gln Arg Leu
 50 55 60
 Gly Leu Pro Thr Ile Arg Ala Arg Val Arg Arg Ala Pro Arg Ser Val
 65 70 75 80
 Leu Asn Leu His Leu Ala
 85

<210> 9
 <211> 87
 <212> PRT
 <213> Nostoc sp.

<400> 9
 Met Val Arg Val Gln Glu Ile Pro Leu Asn Gln Ile Arg Arg Pro Leu
 1 5 10 15

Pro Arg Gly Asn Asp Pro Tyr Lys Val Gln Ala Leu Met Glu Ser Ile
 20 25 30

Ala Ala Ile Gly Gln Gln Glu Pro Ile Asp Val Leu Glu Val Asp Gly
 35 40 45

Gln Tyr Tyr Gly Phe Ser Gly Cys His Arg Tyr Glu Ala Cys Gln Arg
 50 55 60

Leu Gly Lys Glu Thr Ile Leu Ala Arg Val Arg Lys Ala Pro Arg Ser
 65 70 75 80

Val Leu Lys Met His Leu Ala
 85

<210> 10
 <211> 141
 <212> PRT
 <213> Oryza sativa

<400> 10
 Met Ala Ala Ser Gly Phe Leu Leu Arg Cys Pro Ala Ala Pro Ser Ala
 1 5 10 15

Val Pro Leu Trp Gly Arg Ser Gly Arg Gly Gly Gly Gly Gly Leu Ala
 20 25 30

Phe Ser Ala Ser Ser Ser Asn Gly Ala Ala Val Pro Ser Ser Leu Ser
 35 40 45

Asp Ser Glu Lys Lys Gly Pro Val Val Met Glu Ile Pro Leu Asp Lys
 50 55 60

Ile Arg Arg Pro Leu Met Arg Thr Arg Ala Asn Asp Pro Ala Lys Val
 65 70 75 80

Gln Glu Leu Met Asp Ser Ile Arg Val Ile Gly Leu Gln Val Pro Ile
 85 90 95

Asp Val Leu Glu Val Asp Gly Val Tyr Tyr Gly Phe Ser Gly Cys His
 100 105 110

Arg Tyr Glu Ala His Gln Arg Leu Gly Leu Pro Thr Ile Arg Cys Lys
 115 120 125

Val Arg Arg Gly Thr Lys Glu Thr Leu Arg Ile Gly Cys
 130 135 140

<210> 11
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence: Synthetic
 primer

<400> 11
gtcccgcggc ggcggcgacg

20

<210> 12
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 12
agcaggtgcc aaggaggctg

20

<210> 13
<211> 32
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 13
ttaattgaat tcatggggct gcgtgcagga gg

32

<210> 14
<211> 44
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
primer

<400> 14
ttttcctttt gcggccgcct actactgcaa gtctggtgtg gatg

44

<210> 15
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
peptide

<220>
<221> MOD_RES
<222> (2)
<223> Gly or Ser

<400> 15
Phe Xaa Gly Cys His Arg
1 5

<210> 16
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
peptide

<400> 16
Phe Ser Gly Cys His Arg
1 5

<210> 17
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
6xHis tag

<400> 17
His His His His His His
1 5